

Applicants: Bruce D. Gaynor, Betty A. Diamond, Matthew D. Scharff,  
and Philippe Valadon  
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(SEQ ID NO:2), wherein X1 represents Asp or Glu, and X2 represents Gly or Ser; or (iv)  
X1-Gly-X1-Trp-Pro-Arg (SEQ ID NO:5), wherein X1 represents Asp or Glu.

55. (amended) The method according to Claim 54 wherein said  
peptide is 5-30 amino acids in length and comprises X-Gly-Trp-X-Arg-Val (SEQ ID NO:3),  
wherein X represents any amino acid known in the art.

56. (amended) The method according to Claim 54 wherein said  
peptide is 5-15 amino acids in length and comprises X-Gly-Trp-X-Arg-Val (SEQ ID NO:3),  
wherein X represents any amino acid known in the art.

57. (amended) The method according to Claim 54 wherein said  
peptide is 5-10 amino acids in length and comprises X-Gly-Trp-X-Arg-Val (SEQ ID NO:3),  
wherein X represents any amino acid known in the art.

58. (amended) The method according to Claim 54 wherein said  
peptide consists of X-Gly-Trp-X-Arg-Val (SEQ ID NO:3), wherein X represents any amino  
acid known in the art.

59. (amended) The method according to Claim 54 wherein said  
peptide is 5-30 amino acids in length and comprises X-Trp-X-Tyr-His-X (SEQ ID NO:4),  
wherein X represents any amino acid known in the art.

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60. (amended) The method according to Claim 54 wherein said peptide is 5-15 amino acids in length and comprises X-Trp-X-Tyr-His-X (SEQ ID NO:4), wherein X represents any amino acid known in the art.

61. (amended) The method according to Claim 54 wherein said peptide is 5-10 amino acids in length and comprises X-Trp-X-Tyr-His-X (SEQ ID NO:4), wherein X represents any amino acid known in the art.

62. (amended) The method according to Claim 54 wherein said peptide consists of X-Trp-X-Tyr-His-X (SEQ ID NO:4), wherein X represents any amino acid known in the art.

C1  
63. (amended) The method according to Claim 54 wherein said peptide is 5-30 amino acids in length and comprises X1-Trp-X1-Tyr-X2 (SEQ ID NO:2), wherein X1 represents Asp or Glu, and X2 represents Gly or Ser.

64. (amended) The method according to Claim 54 wherein said peptide is 5-15 amino acids in length and comprises X1-Trp-X1-Tyr-X2 (SEQ ID NO:2), wherein X1 represents Asp or Glu, and X2 represents Gly or Ser.

65. (amended) The method according to Claim 54 wherein said peptide is 5-10 amino acids in length and comprises X1-Trp-X1-Tyr-X2 (SEQ ID NO:2), wherein X1 represents Asp or Glu, and X2 represents Gly or Ser.

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C2 66. (amended) The method according to Claim 54 wherein said peptide consists of X1-Trp-X1-Tyr-X2 (SEQ ID NO:2), wherein X1 represents Asp or Glu, and X2 represents Gly or Ser.

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71. (amended) The method according to Claim 54 wherein said peptide is 5-30 amino acids in length and comprises d-Asp-Trp-Glu-Tyr-Ser (SEQ ID NO:2).

Sub D2  
C3 72. (amended) The method according to Claim 54 wherein said peptide is 5-15 amino acids in length and comprises d-Asp-Trp-Glu-Tyr-Ser (SEQ ID NO:2).

73. (amended) The method according to Claim 54 wherein said peptide is 5-10 amino acids in length and comprises d-Asp-Trp-Glu-Tyr-Ser (SEQ ID NO:2).

74. (amended) The method according to Claim 54 wherein said peptide consists of d-Asp-Trp-Glu-Tyr-Ser (SEQ ID NO:2)

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C4 Please attach at the end of the application pages 1-9 of the Sequence Listing (attached hereto as Exhibit B).

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REMARKS

Claims 54-74 were pending in the above-identified application. By this Amendment, applicants have amended the specification to refer to sequence identifiers,